

## SEQUENCE LISTING

<110> Republic of Korea represented by the president of Republic of National Fisheries Research and Development Institute

<120> Phytase produced from *Citrobacter braakii*

<130> 3p-02-25

<160> 8

<170> KopatentIn 1.71

<210> 1

<211> 1481

<212> DNA

<213> *Citrobacter braakii* YH-15

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 <212> PRT  
 <213> *Citrobacter braakii* YH-15

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<210> 3  
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 <212> PRT  
 <213> *Escherichia coli*

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<210> 4  
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 <212> PRT  
 <213> *Aspergillus ficuum*

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 <213> *Bacillus* sp.

<400> 5  
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<210> 6  
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 <212> DNA  
 <213> *Citrobacter braakii* YH-15

<220>  
 <221> gene  
 <222> (1) .. (1302)  
 <223> phytase gene

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<210> 7  
 <211> 433  
 <212> PRT  
 <213> Citrobacter braakii YH-15

<220>  
 <221> PEPTIDE

&lt;222&gt; (1)..(433)

&lt;223&gt; phytase

&lt;400&gt; 7

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Val Val Ile Val Ser Arg His Gly Val Arg Ala Pro Thr Lys Phe Thr
      35           40           45

Pro Ile Met Lys Asp Val Thr Pro Asp Gln Trp Pro Gln Trp Asp Val
 50           55           60

Pro Leu Gly Trp Leu Thr Pro Arg Gly Gly Glu Leu Val Ser Glu Leu
65           70           75           80

Gly Gln Tyr Gln Arg Leu Trp Phe Thr Ser Lys Gly Leu Leu Asn Asn
      85           90           95

Gln Thr Cys Pro Ser Pro Gly Gln Val Ala Val Ile Ala Asp Thr Asp
      100           105           110

Gln Arg Thr Arg Lys Thr Gly Glu Ala Phe Leu Ala Gly Leu Ala Pro
      115           120           125

Lys Cys Gln Ile Gln Val His Tyr Gln Lys Asp Glu Glu Lys Asn Asp
      130           135           140

Pro Leu Phe Asn Pro Val Lys Met Gly Lys Cys Ser Phe Asn Thr Leu
145           150           155           160

Lys Val Lys Asn Ala Ile Leu Glu Arg Ala Gly Gly Asn Ile Glu Leu
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Tyr Thr Gln Arg Tyr Gln Ser Ser Phe Arg Thr Leu Glu Asn Val Leu
      180           185           190

Asn Phe Ser Gln Ser Glu Thr Cys Lys Thr Thr Glu Lys Ser Thr Lys
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Cys Thr Leu Pro Glu Ala Leu Pro Ser Glu Phe Lys Val Thr Pro Asp
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Asn Val Ser Leu Pro Gly Ala Trp Ser Leu Ser Ser Thr Leu Thr Glu
225           230           235           240

Ile Phe Leu Leu Gln Glu Ala Gln Gly Met Pro Gln Val Ala Trp Gly
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Arg Ile Thr Gly Glu Lys Glu Trp Arg Asp Leu Leu Ser Leu His Asn
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Ala Gln Phe Asp Leu Leu Gln Arg Thr Pro Glu Val Ala Arg Ser Arg
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Ala Thr Pro Leu Leu Asp Met Ile Asp Thr Ala Leu Leu Thr Asn Gly  
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Thr Thr Glu Asn Arg Tyr Gly Ile Lys Leu Pro Val Ser Leu Leu Phe  
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Ile Ala Gly His Asp Thr Asn Leu Ala Asn Leu Ser Gly Ala Leu Asp  
 325 330 335

Leu Lys Trp Ser Leu Pro Gly Gln Pro Asp Asn Thr Pro Pro Gly Gly  
 340 345 350

Glu Leu Val Phe Glu Lys Trp Lys Arg Thr Ser Asp Asn Thr Asp Trp  
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Val Gln Val Ser Phe Val Tyr Gln Thr Leu Arg Asp Met Arg Asp Ile  
 370 375 380

Gln Pro Leu Ser Leu Glu Lys Pro Ala Gly Lys Val Asp Leu Lys Leu  
 385 390 395 400

Ile Ala Cys Glu Glu Lys Asn Ser Gln Gly Met Cys Ser Leu Lys Ser  
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Phe Ser Arg Leu Ile Lys Glu Ile Arg Val Pro Glu Cys Ala Val Thr  
 420 425 430

Glu

<210> 8  
 <211> 30  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> primer for the detection of phytase gene

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30